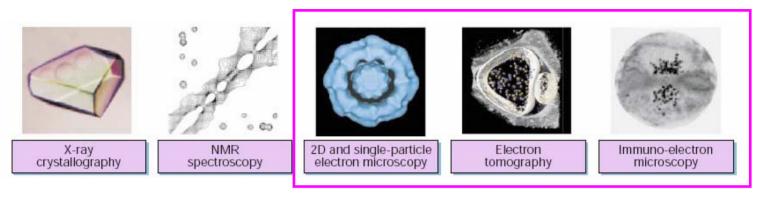
# Cryo electron microscopy

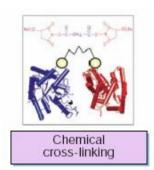
### Capability, Limitation and Potential Solutions

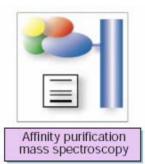
Huilin Li

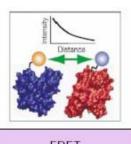
Biology Department Brookhaven National Laboratory

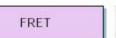
# **Cryo-EM - where does it stand in molecular biology?**

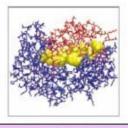




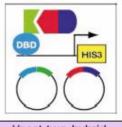












Yeast two-hybrid system

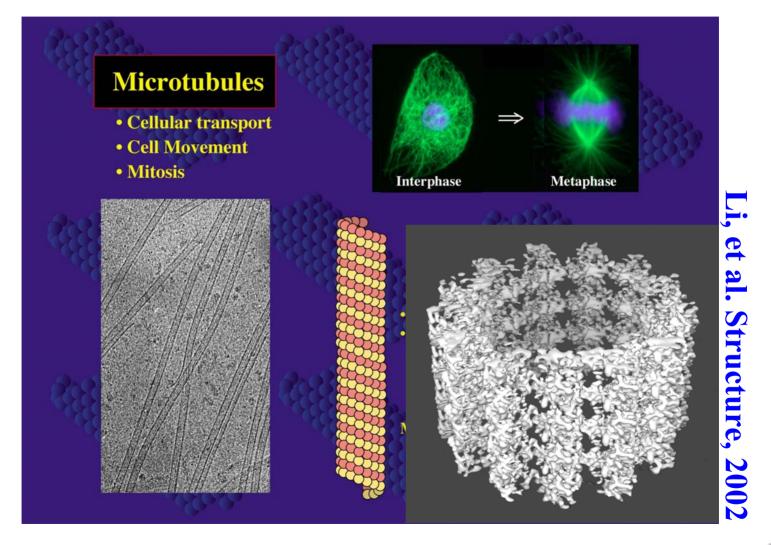
#### Cryo-EM

Samples: purified molecular complexes and microbes

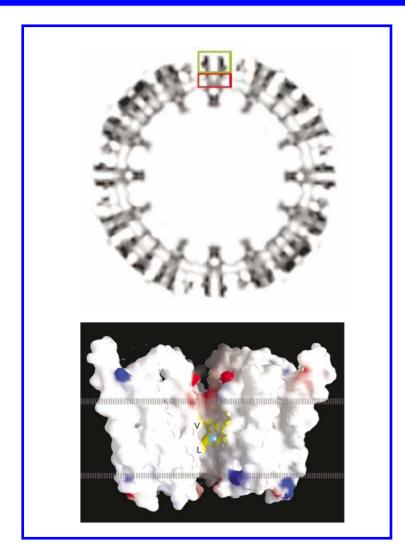
Resolution: 0.35 - 8nm

Biomedical application: structure - function correlation

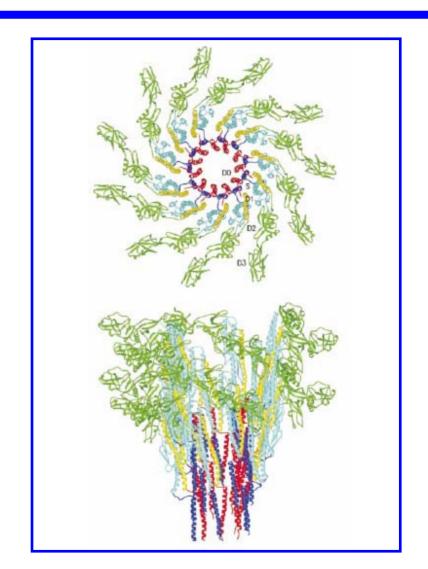
Sample prep: vitrification



#### **Atomic Models of ACh-Receptor and Flagellum by Cryo-EM**



Miyazawa, et al. Nature, 2003



Yonekura, et al. Nature, 2003



Huilin Li Biology

# **Cryo-EM - current limitations**

#### Biggest Problem - 80-90% signal loss

- (a) specimen movement and charging
- (b) modulation transfer function of the film and densitometer
- (c) partial coherence of the electron source
- (d) depth of focus
- (e) beam tilt (f) particle image alignment error

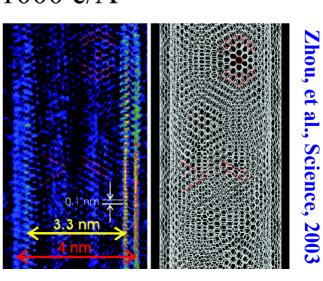
#### Potential solutions

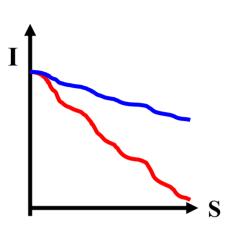
- (1) EM automation and supercomputing
- (2) Diffraction imaging

# CL CA FP OL Sample

# 1000 e/Å<sup>2</sup>

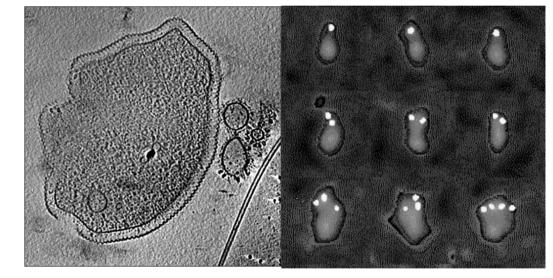
В





## **Cryo-EM - Protein Localization in Microbial Cells**

# Practical problem in imaging microbial cells by tomography molecular crowding



 $\mathsf{QuickTime}^{}{}^{\mathsf{TM}}$  and a TIFF (Uncompressed) decompressor are needed to see this picture.

Potential solution: electron dense labeling

